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Removing near-neighbour **redundancy** from large protein **sequence** collections-

► [oxfordjournals.org](#) (PDF)

L Holm, C Sander - Bioinformatics, 1998 - Oxford Univ Press

... Whenever there is not a **match**, one cannot be certain ... Wootton,JC (1990) Construction of validated, non-**redundant** composite protein **sequence** databases. ...

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The DNA **sequence** of human chromosome 21- ► [fl-leibniz.de](#) (PDF)

M Hattori, A Fujiyama, TD Taylor, H Watanabe ... - Nature, 2000 - nature.com

... of >0.6 and had no **match** to repetitive ... Alu (red) and LINE1 (blue) interspersed repeat element densities ... The final non-**redundant** **sequence** was divided into 340 ...

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TROLL-tandem repeat occurrence locator- ► [oxfordjournals.org](#) (PDF)

AT Castelo, W Martins, GR Gao - Bioinformatics, 2002 - Oxford Univ Press

... m is a **match** for a motif of length i at position k in the DNA **sequence**. ... to account for the appearance of these **redundant** repeats a master record R i is ...

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Rapid, high fidelity analysis of simple **sequence** repeats on an electronically active DNA ...-

► [nih.gov](#) (PDF)

R Radtkey, L Feng, M Muralidhar, M Duhon, D ... - Nucleic Acids Research, 2000 - Oxford Univ Press

... the 1 repeat reporter, the **match** signal appears ... provides the opportunity for **redundant** interrogation of ... interrogate a slightly different **sequence** position of ...

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A map for **sequence** analysis of the Arabidopsis thaliana genome

M Marra, T Kucaba, M Sekhon, LD Hillier, R ... - nature genetics, 1999 - nature.com

... succeed in providing high-quality **sequence**-ready material ... The high **redundancy** achieved by the large number ... best-matching fingerprint.) The best **match** to clone ...

Cited by 94 - Related articles - BL Direct - All 7 versions

[PDF] ► ... RPE/choroid for the NEIBank Project: over 6000 non-**redundant** transcripts, novel genes ...

G Wistow, SL Bernstein, MK Wyatt, RN Fariss, ... - Mol Vis, 2002 - molvis.org

... NEIBank Project: Over 6000 non-**redundant** transcripts, novel ... from lens (by), the RPE/choroid **sequence** data have a ... 45% of the clusters do not **match** named genes ...

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... contains potential coding regions among pseudogenes and a high density of repeat ...- Full

Text@IngentaConnect

M Van Geel, LJ Heather, R Lyle, JE Hewitt, ... - Genomics, 1999 - Elsevier

... clone was initially sequenced with a sixfold **redundancy**. ... A 50-kb **sequence** fragment containing FRG1 and ... predictions usually do not exactly **match** the boundaries ...

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Data mining for simple **sequence** repeats in expressed **sequence** tags from barley, maize, ... -

► [usda.gov](#) (PDF)

RV Kantety, M La Rota, DE Matthews, ME ... - Plant Molecular Biology, 2002 - Springer

... the dataset and creates a longer consensus **sequence** by de ... singletons with a total of 4231 **non-redundant**(NR) **sequences** ... of 85% similarity in order to **match** as many ...

Cited by 250 - Related articles - All 7 versions

► ... **sequence** tag analysis of adult human lens for the NEIBank Project: over 2000 **non-redundant** ...

G Wistow, SL Bernstein, MK Wyatt, A Behal, ... - Mol Vis, 2002 - doaj.org

... NEIBank Project: Over 2000 **non-redundant** transcripts, novel ... the by library contain the complete coding **sequence**. ... the clones have no identifiable **match** with any ...

Cited by 50 - Related articles - Cached - All 5 versions

The complete genome **sequence** of the gastric pathogen *Helicobacter pylori*

JF Tomb, O White, AR Kervavage, RA Clayton, ... - Nature, 1997 - nature.com

... They were searched against a **non-redundant** protein database ... G) tracts occur within the coding **sequence** of nine ... of the PAI have no database **match**, contrary to a ...

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